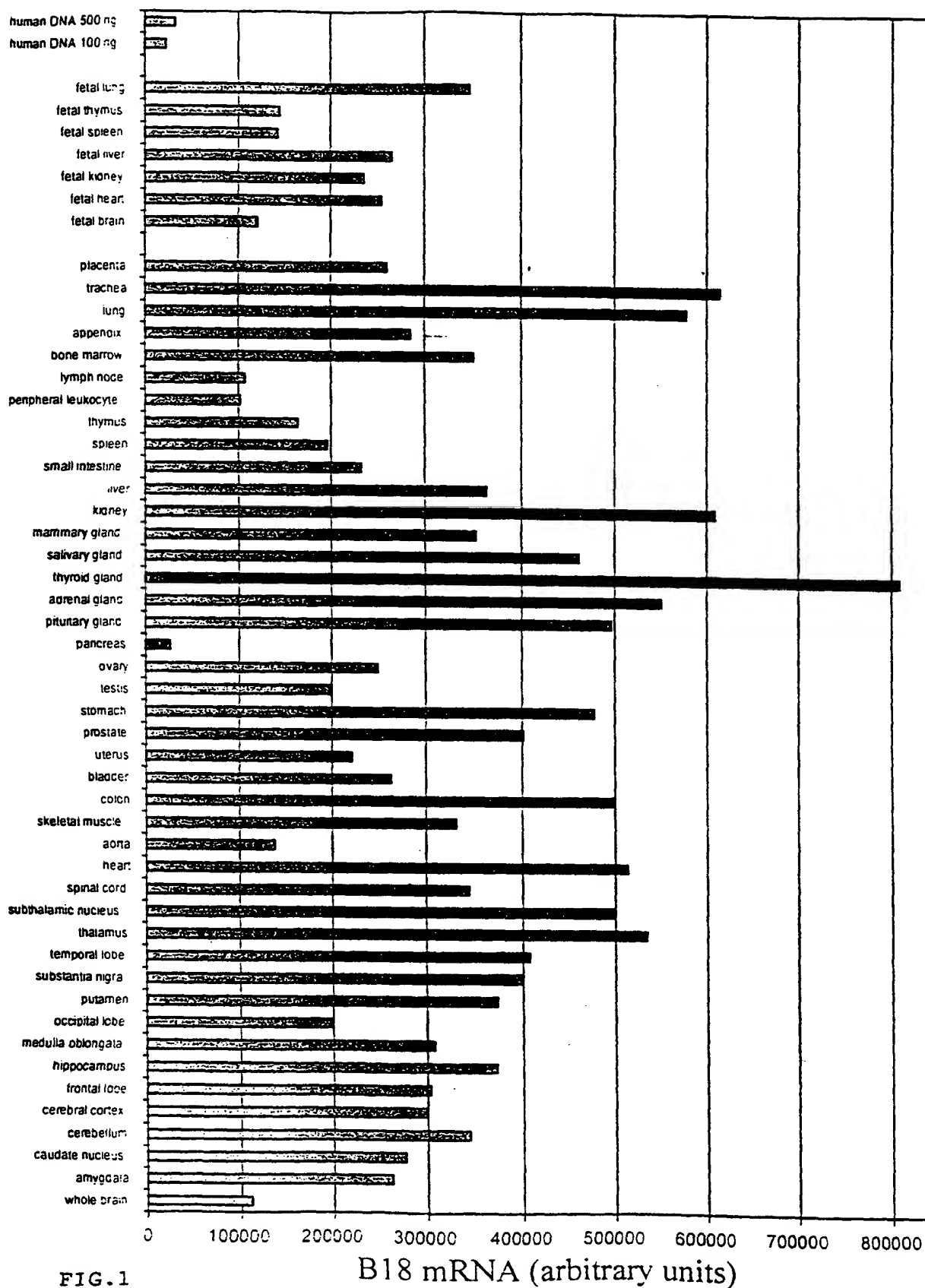
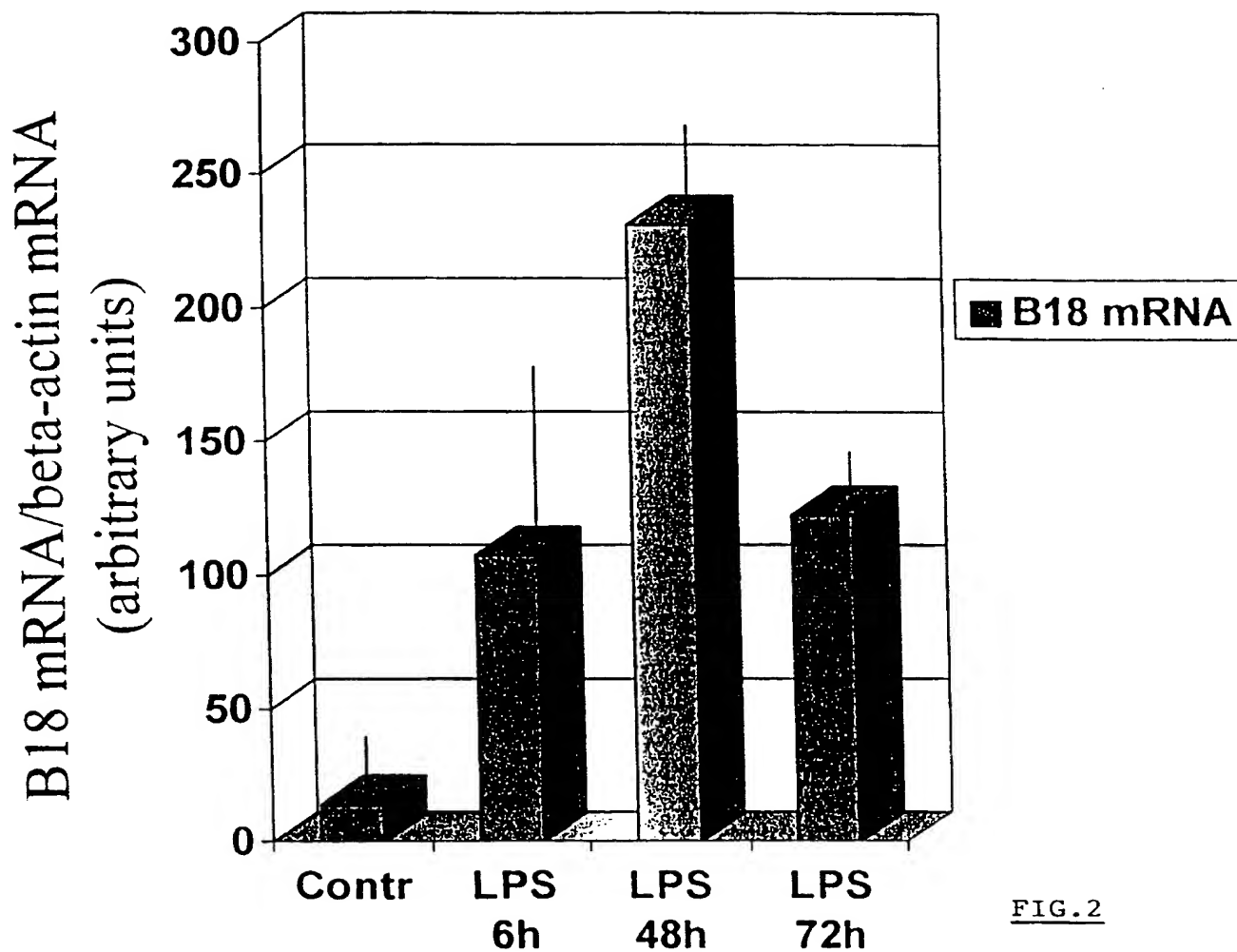


1/7





B18 mRNA/beta-actin mRNA
(arbitrary units)

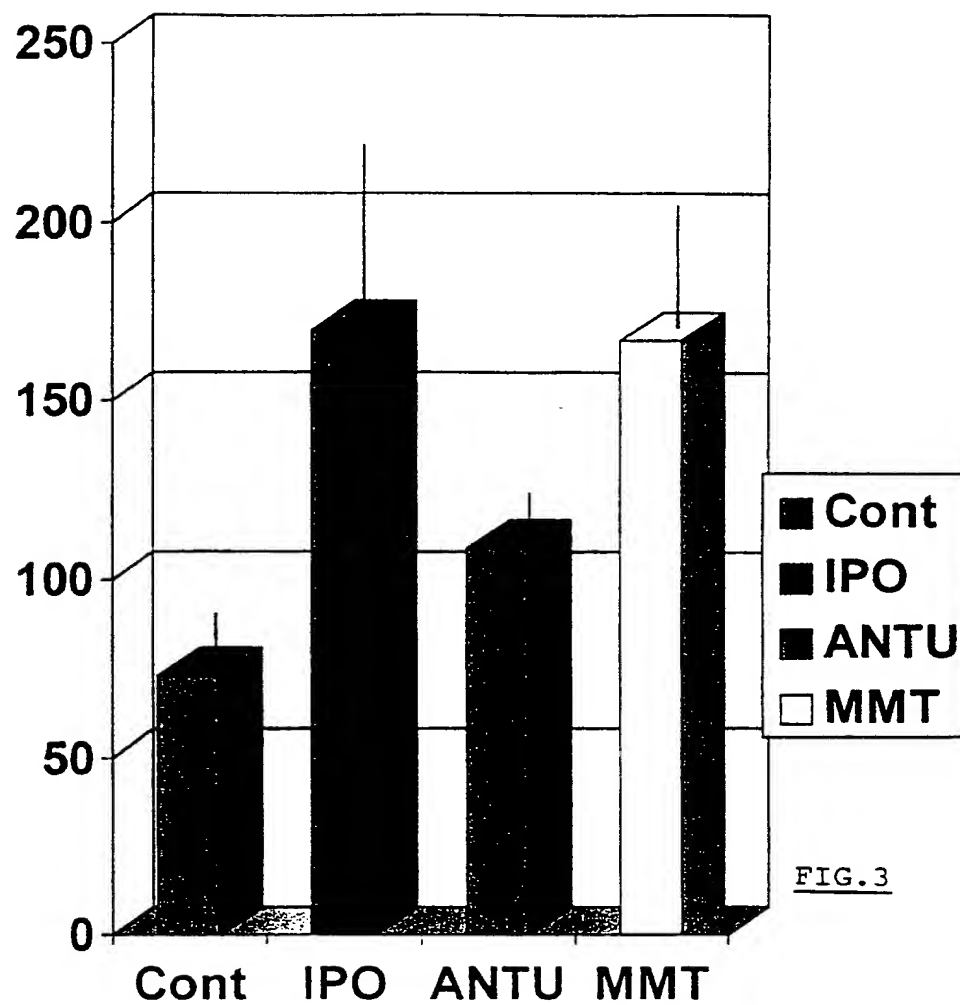
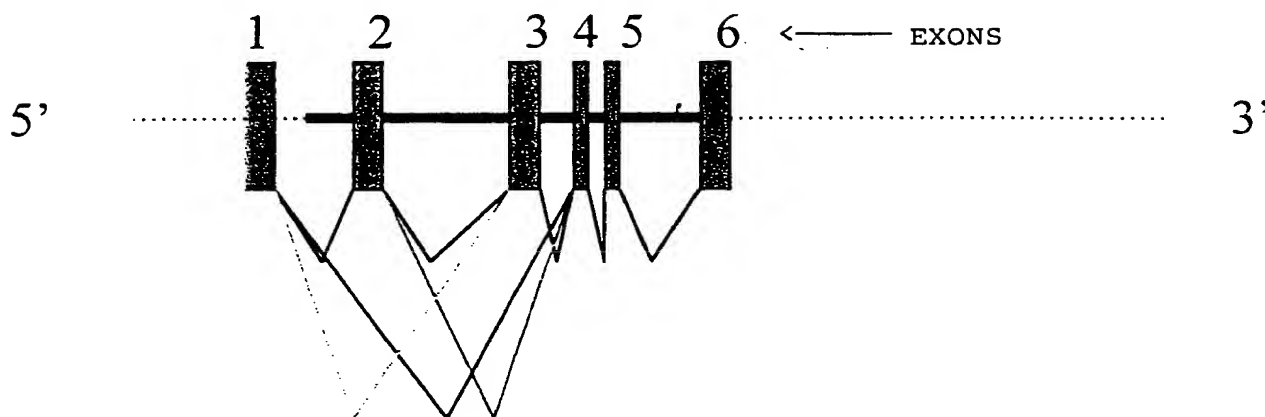
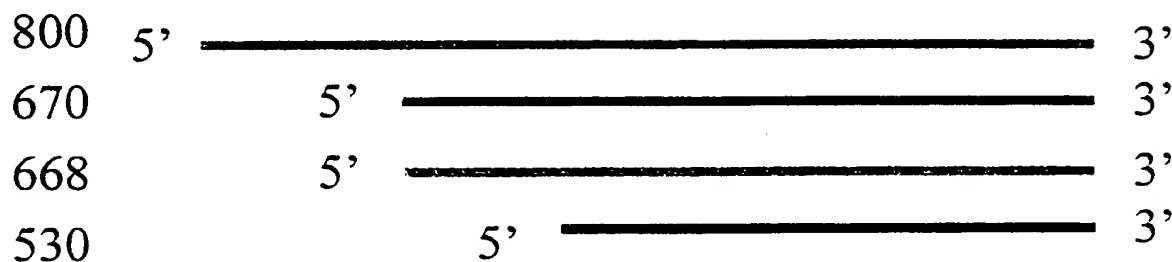


FIG. 3

Gene (chromosome 11q12-13)



mRNAs



Proteins

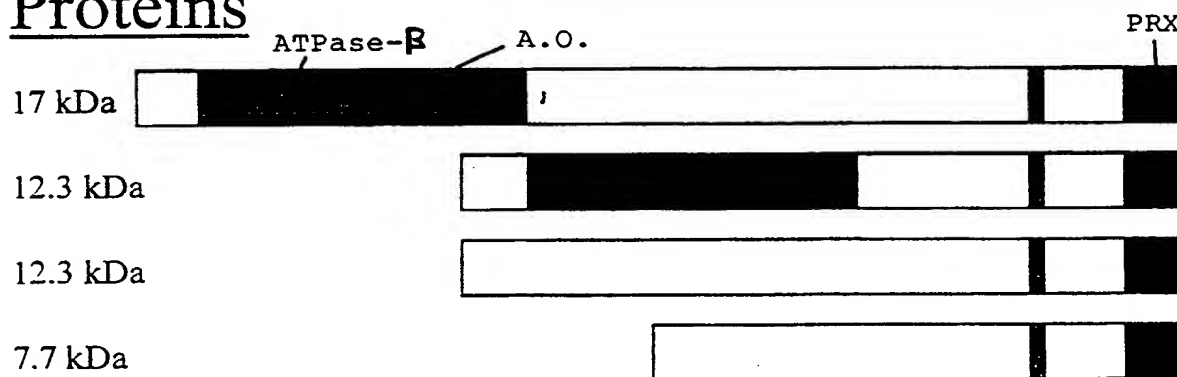


FIG. 4

CLUSTAL V alignment of human and rat B18 amino acid sequences (Identity: 90%, Homology: 97.5%):

```
B10hum      MAPIKVGDAIPAVEVFEGEPGNKVNLAELFKGKKGVLFGVPGAFTPGCSK = SEQIDNO1
B10rat      MAPIKVGDTIPSEVFEGEPGKKVNLAELFKDKKGVLFQVPGAFTPGCSK
*****
```

B18hum THLPGFVEQAEALKAKGVQVVACLSVNDAFVTGEWGRAHKAEGKVRLLD
B18rat THLPGFVEQAGALKAKGAQVVACLSVNDVFVTAEWGRAHQAEQKVQLLD

FIG. 5a

B18hum PTGAFGKETDLLLDDSLVSIFGNRRLKRFSMVVDGIVKALNVEPDGTGL
B18rat PTGAFGKETDLLLDDSLVSLFGNRRLKRFSMVIDKGVVKALNVEPDGTGL
***** * *****

```
B18hum      TCSLAPNIIISQL
B18rat      TCSLAPNIIISQL
            *****
```

CLUSTAL V alignment of human and mouse B18 amino acid sequences (Identity: 91%, Homology: 96%):

B18hum MAPIKVGDAIPAVEVFEGEPGNKVNLAELFKGKKGVLFVGPGAFTPGCSK
B18mouse MAPIKVGDAIPSVEVFEGEPGKKVNLAELFKGKKGVLFVGPGAFTPGCSK

```
B18hum      THLPGFVEQAEALKAKGVQVVACLSVNDVFTGEWGRAHKAEGKVRLLAD
B18mouse    THLPGFVEQAGALKAKGAQVVACLSVNDVFVIEEWGRAHQAEKVRLLAD
*****
```

B18hum PTGAFGKETDLLLDDSLVSI FGNRRLKRFSMVVDGIVKALNVEPDGTGL
B18mouse PTGAFGKATDLLLDDSLVSLFGNRRLKRFSMVIDNGIVKALNVEPDGTGL

```

B18hum      TCSLAPNIIISQL
B18mouse    TCSLAPNILSQL
            *****

```

CLUSTAL V alignment of human and rat cDNA sequences (identity: 612/780, 78.5%):

B18hum GCCAGGAGGCGGAGTGGAAAGTGGCCGTGGGGCGGGTATGGGACTAGCTGG
B18rat -----TG-----CGTC-----CTAGGCAGC

```

B18hum      CGTGTGCGCCCTGAGACGCTCAGCGGGCTATATACTCGTCGGTGGGGCCG
B18rat      CATA---GCC---GGA---TCGGTGCTCCGTGCATCGGCTACTTGGAC--
            * *      ***      **      * * * * *      * *      *      * * * *

```

B18hum GCGGTCAGTCTGCGGCAGCGGCAGCAAGACGGTGCAGTGAAGGAGAGTGG
B18rat -----GTGCGTGGCAGGCAGAGCAGGCCCG--AAAGGAGCAGGTTGG
 * * * * * * * * * * * * * * * * * * * *

6/7

FIG.5b

B18hum GCGTCTGGCGGGGTCCGCAGTTTCAGCAGAGCCGCTGCAGCCATGGCCCC
B18rat GAGTGTGGTGGGGCCCCGCAGCTTCAGCAGTGCCGCGGTGACTATGGCCCC
* * * * *

B18hum AATCAAGGTGGGAGATGCCATCCCAGCAGTGGAGGTGTTTGAAGGGGAGC
B18rat GATCAAGGTGGGAGACACCATTCCCTCAGTGGAGGTATTTGAAGGGGAAC
* * * * *

B18hum CAGGGAACAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAGGGTGTG
B18rat CTGGAAAGAAGGTGAACCTGGCAGAGCTGTTCAAGGACAAGAAAGGTGTT
* * * * *

B18hum CTGTTTGGAGTTCCTGGGGCCTTCACCCCTGGATGTTCCAAGACACACCT
B18rat TTGTTTGGAGTCCCTGGGGCATTACACCTGGCTGTTCCAAGACCCATCT
* * * * *

B18hum GCCAGGGTTTGTGGAGCAGGCTGAGGCTCTGAAGGCCAAGGGAGTCCAGG
B18rat GCCTGGGTTTGTGGAGCAAGCCGGAGCTCTGAAGGCCAAGGGAGCACAAAG
* * * * *

B18hum TGGTGGCCTGTCTGAGTGTTAATGATGCCTTTGTGACTGGCGAGTGGGGC
B18rat TGGTGGCCTGTCTGAGTGTTAATGATGTCTTCGTGACTGCAGAGTGGGGT
* * * * *

B18hum CGAGCCCACAAGGCGGAAGGCAAGGTTTCGGCTCCTGGCTGATCCCCTG
B18rat CGAGCCCACCAGGCAGAAGGCAAGGTTTCAGCTCCTGGCTGACCCCACTGG
* * * * *

B18hum GGCCTTTGGGAAGGAGACAGACTTATTACTAGATGATTTCGCTGGTGTCCA
B18rat AGCTTTTGGAAAGGAGACAGATTTACTACTAGATGATTCTTTGGTGTCTC
* * * * *

B18hum TCTTTGGGAATCGACGTCTCAAGAGGTTCTCCATGGTGGTACAGGATGGC
B18rat TCTTTGGGAATCGTCGGCTAAAAAGGTTCTCCATGGTGTATAGACAAGGGC
* * * * *

B18hum ATAGTGAAGGCCCTGAATGTGGAACCAGATGGCACAGGCCTCACCTGCAG
B18rat GTAGTAAAGGCACTGAACGTGGAGCCGGATGGCACAGGCCTCACCTGCAG
* * * * *

B18hum CCTGGCACCCAATATCATCTCACAGCTCTGAGGCCCTGGGCCAGATTACT
B18rat CCTGGCCCCCAACATCCTCTCACAACTCTGAGGCCCTGA-CCAGA--ATG
* * * * *

B18hum TCCTCCACCCCTCCCTATCTCACCTGCCCAGCCCTGTGCTGG-GGCCCTG
B18rat TCCTCTGACTCTCCC-ATCTCCTCCACCCAGCTCTGGGCCAAAGGCCAG
* * * * *

B18hum CA-----ATTGGAATG-----TTGGCCAGATTTCTGC
B18rat TACCTCCTTACCTGAGGGCCACTGGAATGGAACCTTGACAATATTTCTGC
* * * * *

B18hum AATAAACACTTGTGGTTTGC GGAAAAAA-----
B18rat AATAAACAGTT-TAATTTGTGAAAAAAAAAAAAAAAAAAAA
* * * * *

7/7

CLUSTAL V alignment of human and mouse cDNA sequences (Identity: 552/675, 81.8%):

FIG.5c

```
B18hum      GCCAGGAGGCGGAGTGGAAAGTGGCCGTGGGGCGGGTATGGGACTAGCTGG
B18mouse    -----

B18hum      CGTGTGCGCCCTGAGACGCTCAGCGGGCTATATACTCGTCGGTGGGGCCG
B18mouse    -----TGCTCCGTG-----CATCGACGTGCTTG
                      **** * * * * *

B18hum      GCGGTCAGTCTGCGGCAGCGGCAGCAAGACGGTGCAGTGAAGGAGAGTGG
B18mouse    GCAGGCAG-----AGCAGGCCGG---AAAGAAGCAGGTTGG
                      ** * * * * *

B18hum      GCGTCTGGCGGGGTCCGCAGTTTCAGCAGAGCCGCTGCAGCCATGGCCCC
B18mouse    GAGTGTGGCGGAGCCCGCAGCTTCAGCAGCTCCGCGGTGACCATGGCCCC
                      * * * * *

B18hum      AATCAAGGTGGGAGATGCCATCCCAGCAGTGGAGGTGTTTGAAGGGGAGC
B18mouse    GATCAAGGTGGGAGATGCCATTCCTCAGTGGAGGTATTTGAAGGGGAAC
                      *****

B18hum      CAGGGAACAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAGGGTGTG
B18mouse    CGGGAAAGAAGGTGAACCTGGCAGAGCTGTTCAAGGGCAAGAAAGGTGTT
                      * * * * *

B18hum      CTGTTTGGAGTTCCTGGGGCCTTCACCCCTGGATGTTCCAAGACACACCT
B18mouse    TTGTTTGGAGTCCCTGGGGCATTACACCTGGCTGTTCTAAGACCCACCT
                      *****

B18hum      GCCAGGGTTTGTGGAGCAGGCTGAGGCTCTGAAGGCCAAGGGAGTCCAGG
B18mouse    GCCTGGGTGTTGTGGAGCAAGCTGGAGCTCTGAAGGCTAAGGGAGCGCAGG
                      ***

B18hum      TGGTGGCCTGTCTGAGTGTTAATGATGCCTTTGTGACTGGCGAGTGGGGC
B18mouse    TGGTGGCCTGTCTGAGCGTTAATGACGTCTTTGTGATTGAAGAGTGGGGT
                      *****

B18hum      CGAGCCCACAAGGCGGAAGGCAAGGTTTCGGCTCCTGGCTGATCCCCTGG
B18mouse    CGAGCCCACCAGGCAGAAGGCAAGGTTTCGGCTCCTGGCTGACCCCCTGG
                      *****

B18hum      GGCCTTTGGGAAGGAGACAGACTTATTACTAGATGATTGCTGGTGTCCA
B18mouse    AGCCTTTGGGAAGGCGACAGACTTATTATTGGATGATTCTTTGGTGTCTC
                      *****

B18hum      TCTTTGGGAATCGACGTCTCAAGAGGTTCTCCATGGTGGTACAGGATGGC
B18mouse    TCTTTGGGAATCGTCGGCTGAAAAGGTTCTCCATGGTGATAGACAACGGC
                      *****

B18hum      ATAGTGAAGGCCCTGAATGTGGAACCAGATGGCACAGGCCTCACCTGCAG
B18mouse    ATAGTGAAGGCACTGAACGTGGAGCCAGATGGCACAGGCCTCACCTGCAG
                      *****

B18hum      CCTGGCACCCAATATCATCTCACAGCTCTGAGGCCCTGGGCCAGATTACT
B18mouse    CCTGGCCCCAACATCCTCTCCCAACTCTGAGGCCCTGG-CCAGATG---
                      *****

B18hum      TCCTCCACCCCTCCCTATCTCACCTGCCAGCCCTGTGCTGGGGCCCTGC
B18mouse    TCCTCTGACTCTCCC-ATCTCTCCACCCGGCTCT-----AGGCC----
                      *****

B18hum      AATTGGAATGTTGGCCAGATTTCTGCAATAAACACTTGTGGTTTGCGGAA
B18mouse    ----AAAAGGCTCGGTACCTCCTTACTGGGAGC-CACGT-----
                      * * * * *
```